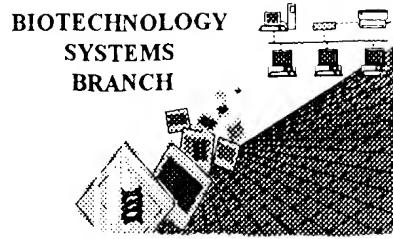


030

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/766,880A
Source: 01P
Date Processed by STIC: 12/13/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

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TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

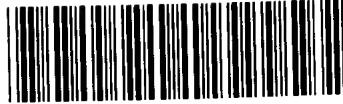
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. **EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)**
2. **U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202**
3. **Hand Carry directly to:**
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
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Revised 01/29/2002

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OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/766,880A

DATE: 12/13/2002
TIME: 15:12:56

Input Set : A:\PTO.txt
Output Set: N:\CRF4\12132002\I766880A.raw

W--> 2 <110> APPLICANT:
W--> 3 Jones, Chris L.
W--> 4 Abrams, Ezra S.
W--> 5 Kieffer-Higgins, Stephen G.
W--> 6 Zhang, Tianhong
W--> 7 Eacock, Graham P.
W--> 8 Webb, Michael
W--> 9 McDowell, Christopher S.
W--> 14 <110> APPLICANT:
W--> 15 Methods of and Apparatus for Separating and Detecting Nucleic Acid
W--> 20 <130> FILE REFERENCE:
W--> 21 2313.2001-001
C--> 26 <140> CURRENT APPLICATION NUMBER:
W--> 27 US 09/766,880
C--> 29 <141> CURRENT FILING DATE:
W--> 30 2001-01-19
W--> 35 <150> PRIOR APPLICATION NUMBER:
W--> 36 US 60/176,839
W--> 38 <151> PRIOR FILING DATE:
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E--> 44 <160> NUMBER OF SEQ ID NOS:
W--> 45 20
50 <170> SOFTWARE:
W--> 51 FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

E--> 56 <210> SEQ ID NO:
W--> 57 1
E--> 59 <211> LENGTH:
W--> 60 50
E--> 62 <212> TYPE:
W--> 63 DNA
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72 1
73 catqaaccat gtcagggtctt gacggaaagca gcattaaatgtt gatccttcata
74 50
E--> 78 <210> SEQ ID NO:
W--> 79 2
E--> 81 <211> LENGTH:

Rec 1.824 of
Sequence Header, all
Sequence Listing files
MUST be saved in
Does Not Comply
Corrected Diskette Needed.
ASCII
Text
format.

The following
pages show
samples of
global errors

FJ1: alphabetical
leading next
to numeric
identifiers were
inserted by CFF

software

If file is created
in MS Word, save it
as MS-DOS Text with Line Breaks

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/766,880A

DATE: 12/13/2002
TIME: 15:12:00

Input Set: A:\PTO.txt
Output Set: N:\CRF4\12132002\I766880A.raw

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E--> 84 <212> TYPE: ~
W--> 85 DNA ~
W--> 87 <213> ORGANISM: ~
W--> 88 unknown
93 <210> FEATURE: ~
96 <213> OTHER INFORMATION: ~
97 Staphylococcus ~
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104 acatgtcagg tcctgacgga agcago ~
105 26 ~
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125 3 ~
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127 50 ~
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W--> 132 4 ~
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W--> 135 49 ~
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156 4 ~
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158 49 ~
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W--> 171 <213> ORGANISM: ~
W--> 172 streptococcus pyogenes
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178 5 ~

4
RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/766,880A

DATE: 12/13/2002

TIME: 10:12:06

Input Set : A:\PTO.txt
Output Set : N:\CRF4\12132002\I766880A.raw

W--> 569 Artificial Sequence

574 <220> FEATURE:

577 <223> OTHER INFORMATION:

578 ~~oligonucleotide probe~~

E--> 583 <400> SEQUENCE:

584 20

585 tccggggccct tgcataaagg agtcccaagg ttccat tggcc g

586 41

589 5/5

592 BOSTON 1546047v1

Delete